logo



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NTDB id 478 HSISS4 RS06985 WP 021143811.1  MTENKNVILSARDIVVEFDVRDRVLTAIRGVSLDLVEGEVLALVGESGSGKSVLTKTFTGMLEENGRVASGSIDYRGQDLTKLKS  85
NTDB id 327 STU RS16115 WP 011226304.1     MTENKNVILSARDIVVEFDVRDRVLTAIRGVSLDLVEGEVLALVGESGSGKSVLTKTFTGMLEENGRVASGSIDYRGKDLTKFKS  85
NTDB id 295 STER RS06915 WP 011226304.1    MTENKNVILSARDIVVEFDVRDRVLTAIRGVSLDLVEGEVLALVGESGSGKSVLTKTFTGMLEENGRVASGSIDYRGKDLTKFKS  85
NTDB id 86369 SI82 RS08940 WP 003102192.1  ..MTTETILNVKNLHVDFKTYAGDVKAIRDLNFELKKGETLAIVGESGSGKSVTTRTLMGLNASNARI.SGDIDFKGKKLNDLK.  81
NTDB id 361 SMU RS01310 WP 002262737.1     ..MSKEKILQVNNLHVNFHTYAGEVKAIRDVSFYLEKGETLAIVGESGSGKSVTTRTLMGLSAKNAEI.FGDIEFKGHNLNDLK.  81
consensus                                  ********!!*******!*!********!!!*****!**!!*!!*!!!!!!!!!!*!*!**!****!*****!*!***!**!***!*
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NTDB id 478 HSISS4 RS06985 WP 021143811.1  HQDWAHIRGAKIATIFQDPMTSLDPIKTIGSQIVEVIVKHQGKSAKEAKKMAIDYMDKVGIPDAEKRFNEYPFQYSGGMRQRIVI  170
NTDB id 327 STU RS16115 WP 011226304.1     HQDWAAIRGAKIATIFQDPMTSLNPIKTIGSQIIEVIVKHQGKTAKEAKKMAIDYMDKVGIPDAEKRFNEYPFQYSGGMRQRIVI  170
NTDB id 295 STER RS06915 WP 011226304.1    HQDWAAIRGAKIATIFQDPMTSLNPIKTIGSQIIEVIVKHQGKTAKEAKKMAIDYMDKVGIPDAEKRFNEYPFQYSGGMRQRIVI  170
NTDB id 86369 SI82 RS08940 WP 003102192.1  EEDWVKVRGKEIAMIFQDPMTSLDPTMKIGMQIAEAILIHEKISKQEALSRALELMHQVGIPNAEEHINDYPHQWSGGMRQRAVI  166
NTDB id 361 SMU RS01310 WP 002262737.1     EEDWVHIRGNDISMIFQDPMTSLDPTMRIGLQIAEPIIKHEKVTKKEALKRALDMMEKVGIPNAQEHINDYPHQWSGGMRQRAVI  166
consensus                                  **!!* *!!**!**!!!!!!!!!*!***!!*!!*!*!**!******!!***!***!**!!!!*!****!*!!*!*!!!!!!!*!!
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NTDB id 478 HSISS4 RS06985 WP 021143811.1  AIALACRPDVLICDEPTTALDVTIQAQIIDLLKSLKEEYGFSVIFITHDLGVVASIADKVAVMYAGEIIEYATVEEIFYEPCHPY  255
NTDB id 327 STU RS16115 WP 011226304.1     AIALACRPDVLICDEPTTALDVTIQAQIIDLLKSLKEEYGFSVIFITHDLGVVASIADKVAVMYAGEIIEYATVEEIFYEPCHPY  255
NTDB id 295 STER RS06915 WP 011226304.1    AIALACRPDVLICDEPTTALDVTIQAQIIDLLKSLKEEYGFSVIFITHDLGVVASIADKVAVMYAGEIIEYATVEEIFYEPCHPY  255
NTDB id 86369 SI82 RS08940 WP 003102192.1  AIALAANPEILIADEPTTALDVTIQNQILKLMKELQSQISSSIIFITHDLGVVAGMADRVAVMYAGKIVEYGTVDEVFYNPQHPY  251
NTDB id 361 SMU RS01310 WP 002262737.1     AIALATNPEILVADEPTTALDVTIQAQILHLMKETQKNTDSSIIFITHDLGVVAGMADRVAVMYAGKIVEYGTVDEVFYNPQHPY  251
consensus                                  !!!!!**!**!**!!!!!!!!!!!!!!*!!**!*!********!*!!!!!!!!!!!!**!!*!!!!!!!*!*!!*!!*!*!!*!*!!!!
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logo



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NTDB id 478 HSISS4 RS06985 WP 021143811.1  TWSLLSSLPQLADDKGELFSIPGTPPSLYTPVVGDAFALRSDYAMQIDFEEKAPQFQVSDTHWAKTWLLHEDAPKVEKPAVIQNL  340
NTDB id 327 STU RS16115 WP 011226304.1     TWSLLSSLPQLADDNGKLFSIPGTPPSLYTPVVGDAFALRSDYALQIDFEEKAPQFQVSDTHWAKTWLLHEDAPKVDKPAVIQNL  340
NTDB id 295 STER RS06915 WP 011226304.1    TWSLLSSLPQLADDNGKLFSIPGTPPSLYTPVVGDAFALRSDYALQIDFEEKAPQFQVSDTHWAKTWLLHEDAPKVDKPAVIQNL  340
NTDB id 86369 SI82 RS08940 WP 003102192.1  TWGLLNSMPTTDTEAGSLESIPGTPPDLLNPPKGDAFAARNEFALDIDHEQEPPMFKVSETHYAATWLLDERAPEVTPPAPILKR  336
NTDB id 361 SMU RS01310 WP 002262737.1     TWGLLNSMPTTNTASGSLHSIPGTPPDLLQPPQGDAFAPRNEFALDIDLKEEPPFFKVSDSHYAATWLLDDRAPKITPPERILQR  336
consensus                                  !!*!!*!*!***** ! !*!!!!!!!*!**!**!!!!!*!***!**!!*****!*!*!!**!*!*!!!!***!!****!**!***
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NTDB id 478 HSISS4 RS06985 WP 021143811.1  HEKIRANMGFAHLGDEEEGNA  361
NTDB id 327 STU RS16115 WP 011226304.1     HEKILANMGFAHLGDEEEGNA  361
NTDB id 295 STER RS06915 WP 011226304.1    HEKILANMGFAHLGDEEEGNA  361
NTDB id 86369 SI82 RS08940 WP 003102192.1  WAKWKKKEGGQA.........  348
NTDB id 361 SMU RS01310 WP 002262737.1     WEKWKSLQGEKHND.......  350
consensus                                  **!******!************
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X non conserved
X similar
X ≥ 50% conserved